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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/662,454

DATE: 09/25/2000  
TIME: 15:16:15

Input Set : A:\20264276.app  
Output Set: N:\CRF3\09252000\I662454.raw

3 <110> APPLICANT: Yanagi, Masayuki  
4 Emerson, Susanne U.  
5 Purcell, Robert H.  
6 Bukh, Jens  
8 <120> TITLE OF INVENTION: CLONED GENOMES OF INFECTIOUS HEPATITIS C VIRUSES AND  
9 USES THEREOF  
11 <130> FILE REFERENCE: 20264276US1  
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/662,454  
C--> 14 <141> CURRENT FILING DATE: 2000-09-14  
16 <150> PRIOR APPLICATION NUMBER: US 60/053,062  
17 <151> PRIOR FILING DATE: 1997-07-18  
19 <150> PRIOR APPLICATION NUMBER: US 09/014,416  
20 <151> PRIOR FILING DATE: 1998-01-27  
22 <160> NUMBER OF SEQ ID NOS: 65  
24 <170> SOFTWARE: PatentIn Ver. 2.1  
26 <210> SEQ ID NO: 1  
27 <211> LENGTH: 3011  
28 <212> TYPE: PRT  
29 <213> ORGANISM: Hepatitis C virus  
31 <400> SEQUENCE: 1  
32 Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn  
33 1 5 10 15  
35 Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly  
36 20 25 30  
38 Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala  
39 35 40 45  
41 Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro  
42 50 55 60  
44 Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly  
45 65 70 75 80  
47 Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp  
48 85 90 95  
50 Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro  
51 100 105 110  
53 Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys  
54 115 120 125  
56 Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu  
57 130 135 140  
59 Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp  
60 145 150 155 160  
62 Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile  
63 165 170 175  
65 Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr  
66 180 185 190  
68 Gln Val Arg Asn Ser Ser Gly Leu Tyr His Val Thr Asn Asp Cys Pro  
69 195 200 205  
71 Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Ala Ile Leu His Thr Pro

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72	210	215	220
74	Gly Cys Val Pro Cys Val Arg Glu Gly Asn Ala Ser Arg Cys Trp Val		
75	225	230	235
			240
77	Ala Val Thr Pro Thr Val Ala Thr Arg Asp Gly Lys Leu Pro Thr Thr		
78	245	250	255
80	Gln Leu Arg Arg His Ile Asp Leu Leu Val Gly Ser Ala Thr Leu Cys		
81	260	265	270
83	Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val Gly		
84	275	280	285
86	Gln Leu Phe Thr Phe Ser Pro Arg Arg His Trp Thr Thr Gin Asp Cys		
87	290	295	300
89	Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met Ala Trp		
90	305	310	315
			320
92	Asp Met Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ala Gln		
93	325	330	335
95	Leu Leu Arg Ile Pro Gln Ala Ile Met Asp Met Ile Ala Gly Ala His		
96	340	345	350
98	Trp Gly Val Leu Ala Gly Ile Ala Tyr Phe Ser Met Val Gly Asn Trp		
99	355	360	365
101	Ala Lys Val Leu Val Val Leu Leu Leu Phe Ala Gly Val Asp Ala Glu		
102	370	375	380
104	Thr His Val Thr Gly Gly Asn Ala Gly Arg Thr Thr Ala Gly Leu Val		
105	385	390	395
			400
107	Gly Leu Leu Thr Pro Gly Ala Lys Gln Asn Ile Gln Leu Ile Asn Thr		
108	405	410	415
110	Asn Gly Ser Trp His Ile Asn Ser Thr Ala Leu Asn Cys Asn Glu Ser		
111	420	425	430
113	Leu Asn Thr Gly Trp Leu Ala Gly Leu Phe Tyr Gln His Lys Phe Asn		
114	435	440	445
116	Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser Cys Arg Arg Leu Thr Asp		
117	450	455	460
119	Phe Ala Gln Gly Trp Gly Pro Ile Ser Tyr Ala Asn Gly Ser Gly Leu		
120	465	470	475
			480
122	Asp Glu Arg Pro Tyr Cys Trp His Tyr Pro Pro Arg Pro Cys Gly Ile		
123	485	490	495
125	Val Pro Ala Lys Ser Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser		
126	500	505	510
128	Pro Val Val Val Gly Thr Thr Asp Arg Ser Gly Ala Pro Thr Tyr Ser		
129	515	520	525
131	Trp Gly Ala Asn Asp Thr Asp Val Phe Val Leu Asn Asn Thr Arg Pro		
132	530	535	540
134	Pro Leu Gly Asn Trp Phe Gly Cys Thr Trp Met Asn Ser Thr Gly Phe		
135	545	550	555
			560
137	Thr Lys Val Cys Gly Ala Pro Pro Cys Val Ile Gly Gly Val Gly Asn		
138	565	570	575
140	Asn Thr Leu Leu Cys Pro Thr Asp Cys Phe Arg Lys His Pro Glu Ala		
141	580	585	590
143	Thr Tyr Ser Arg Cys Gly Ser Gly Pro Trp Ile Thr Pro Arg Cys Met		
144	595	600	605

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146 Val Asp Tyr Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Ile Asn Tyr  
147 610 615 620  
149 Thr Ile Phe Lys Val Arg Met Tyr Val Gly Gly Val Glu His Arg Leu  
150 625 630 635 640  
152 Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu Arg Cys Asp Leu Glu Asp  
153 645 650 655  
155 Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu Ser Thr Thr Gln Trp  
156 660 665 670  
158 Gln Val Leu Pro Cys Ser Phe Thr Thr Leu Pro Ala Leu Ser Thr Gly  
159 675 680 685  
161 Leu Ile His Leu His Gln Asn Ile Val Asp Val Gln Tyr Leu Tyr Gly  
162 690 695 700  
164 Val Gly Ser Ser Ile Ala Ser Trp Ala Ile Lys Trp Glu Tyr Val Val  
165 705 710 715 720  
167 Leu Leu Phe Leu Leu Ala Asp Ala Arg Val Cys Ser Cys Leu Trp  
168 725 730 735  
170 Met Met Leu Leu Ile Ser Gln Ala Glu Ala Ala Leu Glu Asn Leu Val  
171 740 745 750  
173 Ile Leu Asn Ala Ala Ser Leu Ala Gly Thr His Gly Leu Val Ser Phe  
174 755 760 765  
176 Leu Val Phe Phe Cys Phe Ala Trp Tyr Leu Lys Gly Arg Trp Val Pro  
177 770 775 780  
179 Gly Ala Val Tyr Ala Leu Tyr Gly Met Trp Pro Leu Leu Leu Leu  
180 785 790 795 800  
182 Leu Ala Leu Pro Gln Arg Ala Tyr Ala Leu Asp Thr Glu Val Ala Ala  
183 805 810 815  
185 Ser Cys Gly Gly Val Val Leu Val Gly Leu Met Ala Leu Thr Leu Ser  
186 820 825 830  
188 Pro Tyr Tyr Lys Arg Tyr Ile Ser Trp Cys Met Trp Trp Leu Gln Tyr  
189 835 840 845  
191 Phe Leu Thr Arg Val Glu Ala Gln Leu His Val Trp Val Pro Pro Leu  
192 850 855 860  
194 Asn Val Arg Gly Gly Arg Asp Ala Val Ile Leu Leu Met Cys Val Val  
195 865 870 875 880  
197 His Pro Thr Leu Val Phe Asp Ile Thr Lys Leu Leu Ala Ile Phe  
198 885 890 895  
200 Gly Pro Leu Trp Ile Leu Gln Ala Ser Leu Leu Lys Val Pro Tyr Phe  
201 900 905 910  
203 Val Arg Val Gln Gly Leu Leu Arg Ile Cys Ala Leu Ala Arg Lys Ile  
204 915 920 925  
206 Ala Gly Gly His Tyr Val Gln Met Ala Ile Ile Lys Leu Gly Ala Leu  
207 930 935 940  
209 Thr Gly Thr Tyr Val Tyr Asn His Leu Thr Pro Leu Arg Asp Trp Ala  
210 945 950 955 960  
212 His Asn Gly Leu Arg Asp Leu Ala Val Ala Val Glu Pro Val Val Phe  
213 965 970 975  
215 Ser Arg Met Glu Thr Lys Leu Ile Thr Trp Gly Ala Asp Thr Ala Ala  
216 980 985 990  
218 Cys Gly Asp Ile Ile Asn Gly Leu Pro Val Ser Ala Arg Arg Gly Gln

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219	995	1000	1005
221	Glu Ile Leu Leu Gly Pro Ala Asp Gly Met Val Ser Lys Gly Trp Arg		
222	1010	1015	1020
224	Leu Leu Ala Pro Ile Thr Ala Tyr Ala Gln Gln Thr Arg Gly Leu Leu		
225	1025	1030	1035
227	Gly Cys Ile Ile Thr Ser Leu Thr Gly Arg Asp Lys Asn Gln Val Glu		1040
228	1045	1050	1055
230	Gly Glu Val Gln Ile Val Ser Thr Ala Thr Gln Thr Phe Leu Ala Thr		
231	1060	1065	1070
233	Cys Ile Asn Gly Val Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg		
234	1075	1080	1085
236	Thr Ile Ala Ser Pro Lys Gly Pro Val Ile Gln Met Tyr Thr Asn Val		
237	1090	1095	1100
239	Asp Gln Asp Leu Val Gly Trp Pro Ala Pro Gln Gly Ser Arg Ser Leu		
240	1105	1110	1115
242	1120		
242	Thr Pro Cys Thr Cys Gly Ser Ser Asp Leu Tyr Leu Val Thr Arg His		
243	1125	1130	1135
245	Ala Asp Val Ile Pro Val Arg Arg Gly Asp Ser Arg Gly Ser Leu		
246	1140	1145	1150
248	Leu Ser Pro Arg Pro Ile Ser Tyr Leu Lys Gly Ser Ser Gly Gly Pro		
249	1155	1160	1165
251	Leu Leu Cys Pro Ala Gly His Ala Val Gly Leu Phe Arg Ala Ala Val		
252	1170	1175	1180
254	Cys Thr Arg Gly Val Ala Lys Ala Val Asp Phe Ile Pro Val Glu Asn		
255	1185	1190	1195
257	1200		
257	Leu Gly Thr Thr Met Arg Ser Pro Val Phe Thr Asp Asn Ser Ser Pro		
258	1205	1210	1215
260	Pro Ala Val Pro Gln Ser Phe Gln Val Ala His Leu His Ala Pro Thr		
261	1220	1225	1230
263	Gly Ser Gly Lys Ser Thr Lys Val Pro Ala Ala Tyr Ala Ala Gln Gly		
264	1235	1240	1245
266	Tyr Lys Val Leu Val Leu Asn Pro Ser Val Ala Ala Thr Leu Gly Phe		
267	1250	1255	1260
269	Gly Ala Tyr Met Ser Lys Ala His Gly Val Asp Pro Asn Ile Arg Thr		
270	1265	1270	1275
272	1280		
272	Gly Val Arg Thr Ile Thr Thr Gly Ser Pro Ile Thr Tyr Ser Thr Tyr		
273	1285	1290	1295
275	Gly Lys Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Ile		
276	1300	1305	1310
278	Ile Ile Cys Asp Glu Cys His Ser Thr Asp Ala Thr Ser Ile Leu Gly		
279	1315	1320	1325
281	Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val		
282	1330	1335	1340
284	Val Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Val Ser His Pro		
285	1345	1350	1355
287	1360		
287	Asn Ile Glu Glu Val Ala Leu Ser Thr Thr Gly Glu Ile Pro Phe Tyr		
288	1365	1370	1375
290	Gly Lys Ala Ile Pro Leu Glu Val Ile Lys Gly Gly Arg His Leu Ile		
291	1380	1385	1390

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293 Phe Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala Ala Lys Leu Val  
 294 1395 1400 1405  
 296 Ala Leu Gly Ile Asn' Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ser  
 297 1410 1415 1420  
 299 Val Ile Pro Thr Ser Gly Asp Val Val Val Val Ser Thr Asp Ala Leu  
 300 1425 1430 1435 1440  
 302 Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Thr  
 303 1445 1450 1455  
 305 Cys Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile  
 306 1460 1465 1470  
 308 Glu Thr Thr Thr Leu Pro Gln Asp Ala Val Ser Arg Thr Gln Arg Arg  
 309 1475 1480 1485  
 311 Gly Arg Thr Gly Arg Gly Lys Pro Gly Ile Tyr Arg Phe Val Ala Pro  
 312 1490 1495 1500  
 314 Gly Glu Arg Pro Ser Gly Met Phe Asp Ser Ser Val Leu Cys Glu Cys  
 315 1505 1510 1515 1520  
 317 Tyr Asp Ala Gly Cys Ala Trp Tyr Glu Leu Thr Pro Ala Glu Thr Thr  
 318 1525 1530 1535  
 320 Val Arg Leu Arg Ala Tyr Met Asn Thr Pro Gly Leu Pro Val Cys Gln  
 321 1540 1545 1550  
 323 Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr His Ile  
 324 1555 1560 1565  
 326 Asp Ala His Phe Leu Ser Gln Thr Lys Gln Ser Gly Glu Asn Phe Pro  
 327 1570 1575 1580  
 329 Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Gln Ala Pro  
 330 1585 1590 1595 1600  
 332 Pro Pro Ser Trp Asp Gln Met Trp Lys Cys Leu Ile Arg Leu Lys Pro  
 333 1605 1610 1615  
 335 Thr Leu His Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Ala Val Gln  
 336 1620 1625 1630  
 338 Asn Glu Val Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Thr Cys  
 339 1635 1640 1645  
 341 Met Ser Ala Asp Leu Glu Val Val Thr Ser Thr Trp Val Leu Val Gly  
 342 1650 1655 1660  
 344 Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Thr Gly Cys Val  
 345 1665 1670 1675 1680  
 347 Val Ile Val Gly Arg Ile Val Leu Ser Gly Lys Pro Ala Ile Ile Pro  
 348 1685 1690 1695  
 350 Asp Arg Glu Val Leu Tyr Gln Glu Phe Asp Glu Met Glu Glu Cys Ser  
 351 1700 1705 1710  
 353 Gln His Leu Pro Tyr Ile Glu Gln Gly Met Met Leu Ala Glu Gln Phe  
 354 1715 1720 1725  
 356 Lys Gln Lys Ala Leu Gly Leu Leu Gln Thr Ala Ser Arg His Ala Glu  
 357 1730 1735 1740  
 359 Val Ile Thr Pro Ala Val Gln Thr Asn Trp Gln Lys Leu Glu Val Phe  
 360 1745 1750 1755 1760  
 362 Trp Ala Lys His Met Trp Asn Phe Ile Ser Gly Ile Gln Tyr Leu Ala  
 363 1765 1770 1775  
 365 Gly Leu Ser Thr Leu Pro Gly Asn Pro Ala Ile Ala Ser Leu Met Ala

VERIFICATION SUMMARY  
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L:13 M:270 C: Current Application Number differs, Replaced Application Number  
L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date